#7

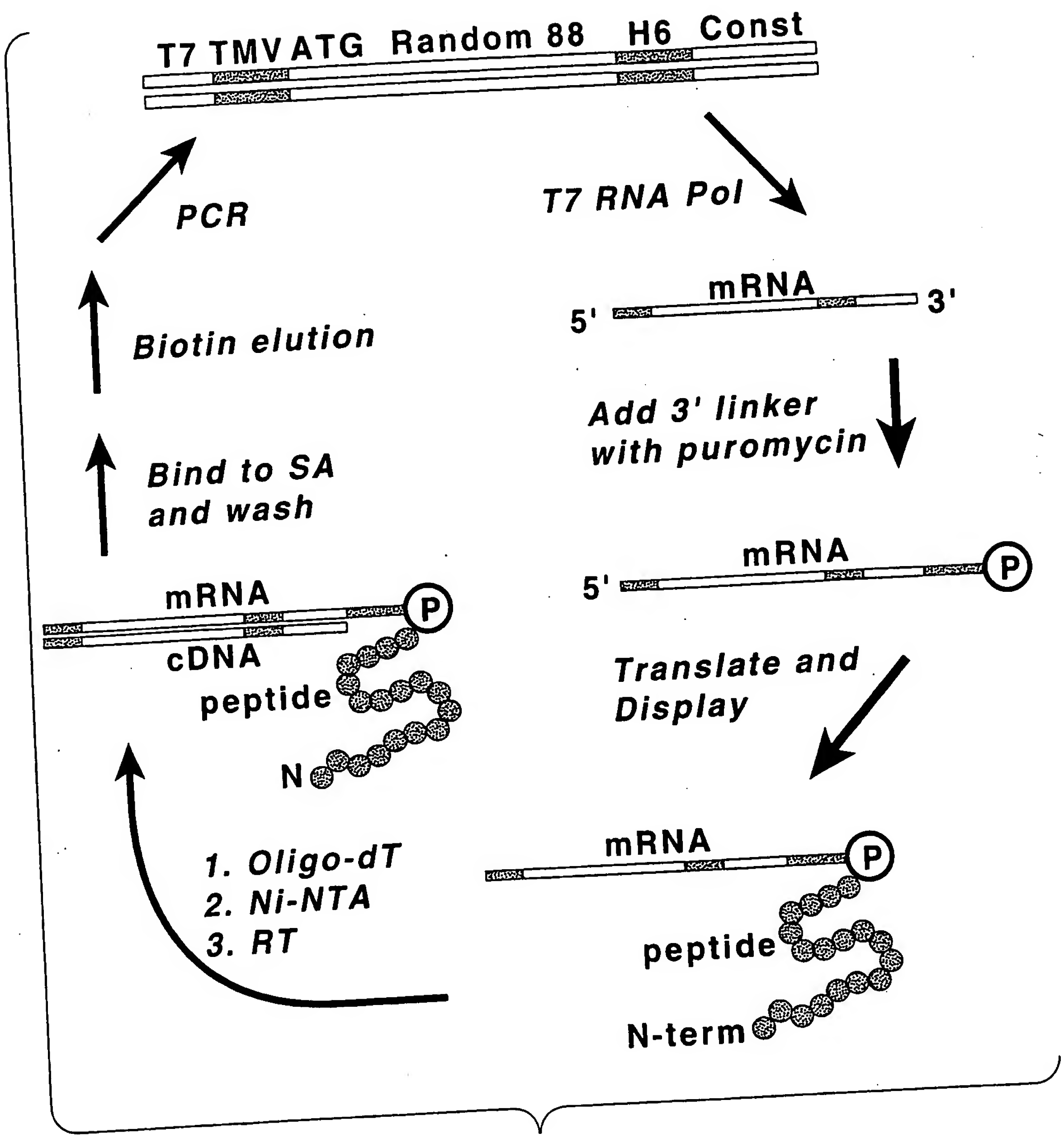

Fig. 1A

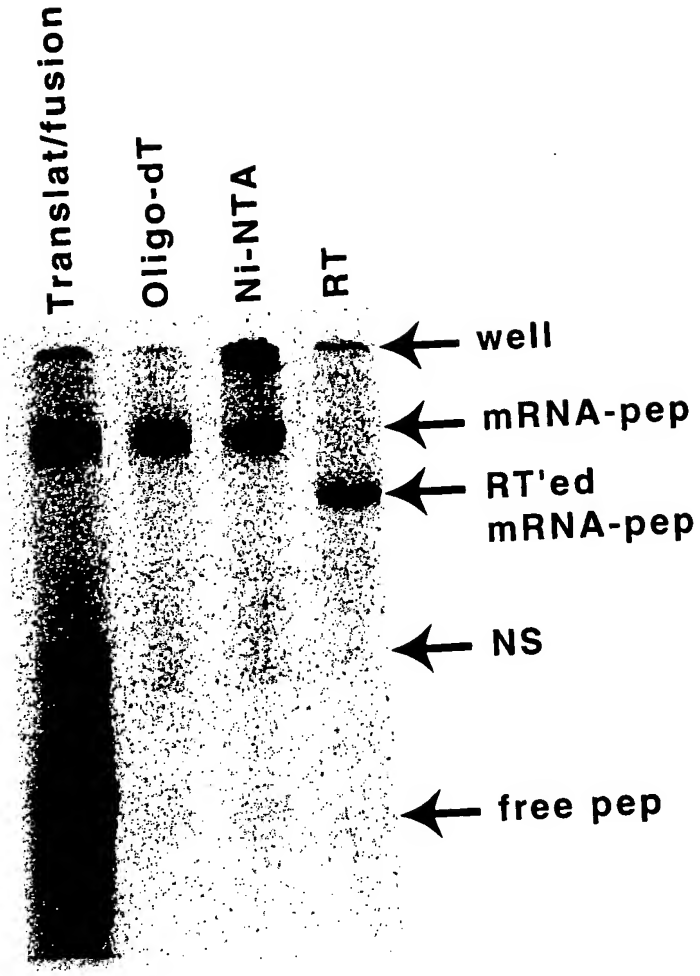

Fig. 1B

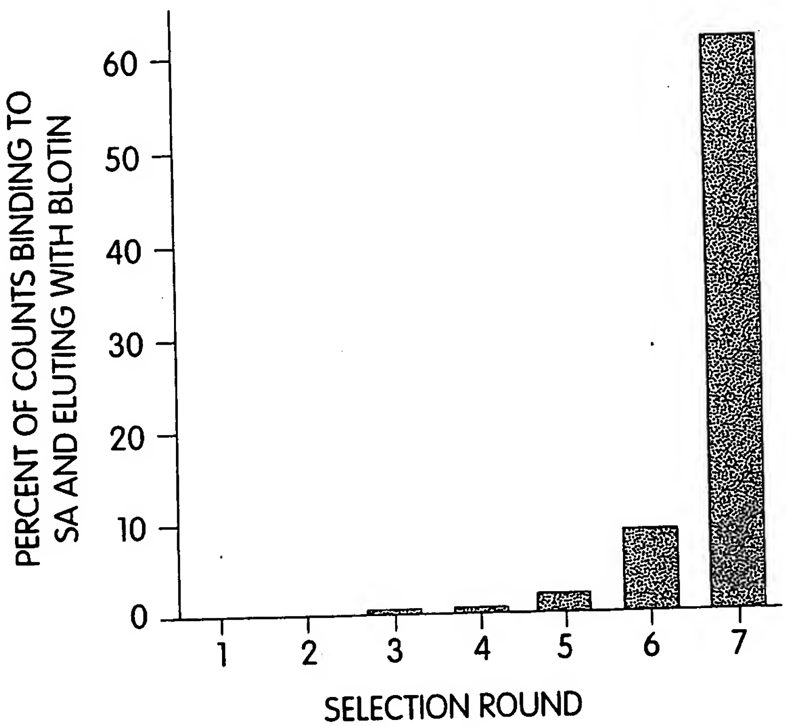
PERCENT OF COUNTS BINDING TO SA AND ELUTING WITH BIOTIN
60
50
40
30
20
10
0
1
2
3
4
5
6
7
SELECTION ROUND

Fig. 2A

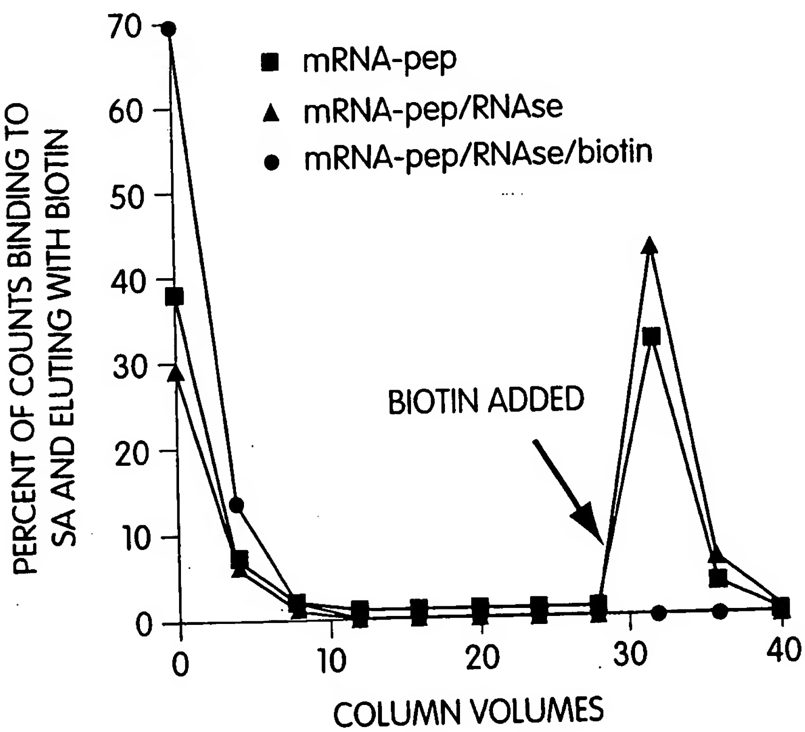
PERCENT OF COUNTS BINDING TO SA AND ELUTING WITH BIOTIN
70
60
50
40
30
20
10
0
mRNA-pep
mRNA-pep/RNAse
mRNA-pep/RNAse/biotin
BIOTIN ADDED
0
10
20
30
40
COLUMN VOLUMES

Fig. 2B

name	#		SEQ ID NO:
SB1	3	MDEKTHCTISMNGAVPLVPHH**HPQ**GDPLRLLHRPQPALLVR**HPQ**GDLVALVEHHEGVDRGLVALPELHAEELGEPVGDLVQGPVEQVQGVVDALVWRLPPS	1
SB2	2	MDEKTHCFHPGDHLVRLVEELQALAEGLQRQGGRQPHRLPRRRPHHLQLLLDEA**HPQ**AGPLRERAHQVDGRLLLQH**HPQ**GDRLLQQPQDHPLELVWRLPPS	2
SB3	4	MTRRPTASSSSCVRHLLLRQGEHGHQALEDRDKARHVRLVEGDVEVLGGLDRLARARHEAL**HPQ**AGLVHLPLHGGDLGGHLRLVLEA**HPQ**GDRLGLAVHHH	3
SB4	1	MDEKTHWGISTWRGEPLLH**HPQ**AGRLPLDRRRARHRRILGAEPGGVDHGLRLELLDDHRPLVPDH**HPQ**RGPLQRGDLPQVVPLVRLRHAHVLGLGLAAATIT	4
SB5	3	MDEKTHWVNVY**HPQ**GDLLVRGHGHDVEALHDQGLHQLDLLVGPPPEVVRALRGEVLGGLRRLVPLD**HPQ**GEALDQARQRPQHLLELHHRALPPALVWRLPPS	5
SB6	1	MDEKTHWLNNFEELLARLDGLREGEDHPLVLRH**HPQ**GDGLLDQPLGRHRALDGEVREGDRPLDQGGEEDLGALVDDDGEVLDGLVHVGVHVHDPLVCGCHHH	6
SB7	1	MDEKTHWFGTLNSFPTHWMSAVGNGKIDCSFNMNLSLNHWLSSGHPDGALDDQL**HPQ**GDALVGRDDGVVQALRLEGQHQHRRLAQRRADRHRQLVWRLPPS	7
SB8	1	MDEKTHCTIELNFSFTHWKLHH**HPQ**GDALLDDGVRPHHPLADEGGGLDQGLGHRRGVVAERLARRDPEVLEGLVERHRGLVPRLRHGGERHAEPLVWRLPPS	8
SB9	1	MDEKTHCNTGLYDGAADCFNELNKDVAPLVEGRHDLVEGLLLER**HPQ**GDPLVAHRQLVHHPLLGRGERHRRALVPQQEHQPHRLQPVVDLGRRRLVWRLPPS	9
SB10	1	MDEKTHWHERAQELVGGLLLHD**HPQ**RLLLEPRGPRPLRGLVHERGHQPQPLAGRVEEADGGLLRDGGGELEPLVREGEDHLEPLDDELDAGPRGLVWRLPHHH	10
SB11	1	MDEKTHWHERVHHLADGLEQ**HPQ**GQRRPLVERHRQVPRGLVRELQHEGLPLEHPAGVHVIRLHQGDDRDVDGLVDGHGRDVRGLEREVGDGPHRLVWRLPPS	11
SB12	4	MDKDPLLEELEELRERLVH**HPQ**GGLLPLRGQVGHDAERLGAEVDDLRGGLLDEPQRAVAGLHHVPHRVGQRLVHEVRELDEGLLDQRDDLRQRLVWRLPPS	12
SB13	2	MEREDPLDEQLRELREALVD**HPQ**GGAQALHRHDGGEHVPLRRVQHRLQPGLQHHLEPQPLGLLGELQARLQPLAGEHEGDGAGLQRVPGHQGRRLVWRLPPS	13
SB14	1	MDEKTHRTLSVSLSFNDWLGQTKACWRLVEGLHG**HPQ**GLVREHEVDVLPLAEEVQQVVGGLADGVEQPGGGLLHRAQRVDHPLPDHAGQVLGRLVWRLPPS	14
SB15	1	MDEKTHWLEDLKGVLKDCLKDLMDFTKDCRSPRVQPQPLLHHDRGEPVPLLREAGRDLGGLGPRAPRQARPLHHGRHDLHEPLVLQD**HPQ**GGPLVCGCHHH	15
SB16	1	MDEKTHWVLQL**HPQ**GDRLGPRHGGDDVRLVGQGEGVLEGLDGRPRRRRHRLPREDEHRVRALVDQVRDLAERLVEEVDGGVEALRHLGLPQDEPRSGGCHHH	16
SB17	2	MDEKTHWVGDLQEPLGPLHGGVGEVPGGLVLRH**HPQ**RDRLVDGVGPHGRALARRPHRVVEGLHHLLQRGGERLPPDGPRQLGLLGGELDRADPALVWRLPPS	17
SB18	1	MDEKTHCAVNVNVGLTHWCHRVAHLQPLDP**HPQ**GDHLRLEPLGHALVDPLVQGVEEVVRPLQLDVGVQRVALVEQVAEVGEGLDHEAGQAHGALVWRLPPS	18
SB19	1	MDEKTTGWRGGHVVEGLAGELEQLRARLEH**HPQ**GQREPLVQEVEDVDEGLVQDLHGVVAGLLDPVEKLLTDWFKKFKNVSKDCKMTFYLEMYDWSGGCHHH	19
SB20	1	MNEKTHCKLNFKVNIADWLAEFHGGGQGLLGRRDGVVQRLVDGVQERVERLDRDPGLGDLRLELHHRDHRLRLGGEHLLRDHPLEPDDHLVVGGLVWRLPPS	20

Fig. 3

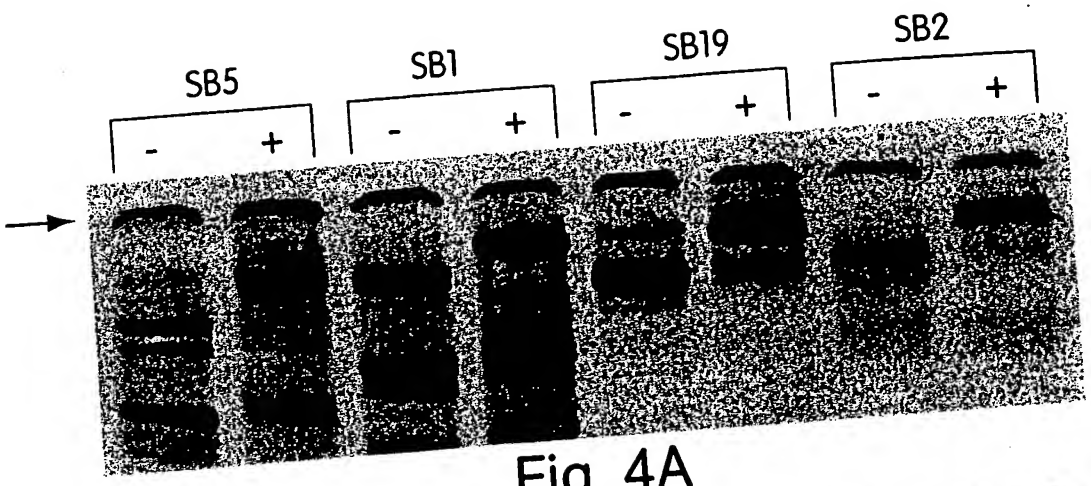

Fig. 4A

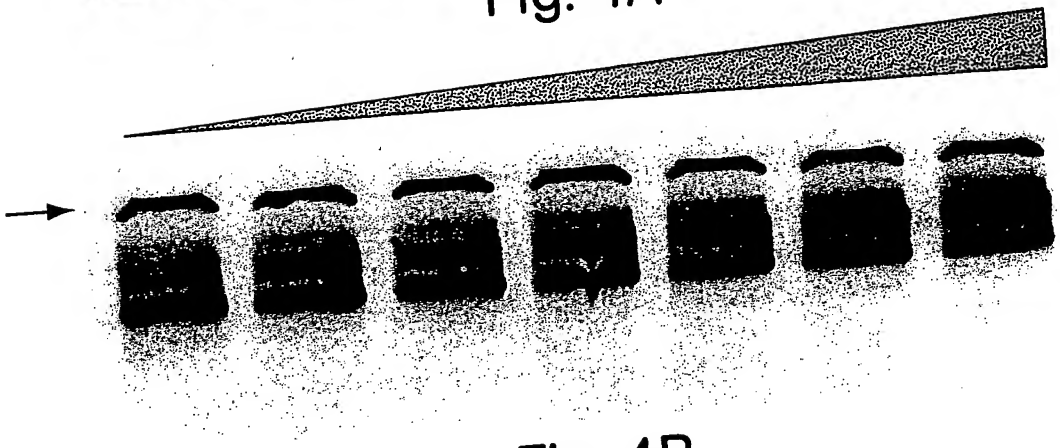

Fig. 4B

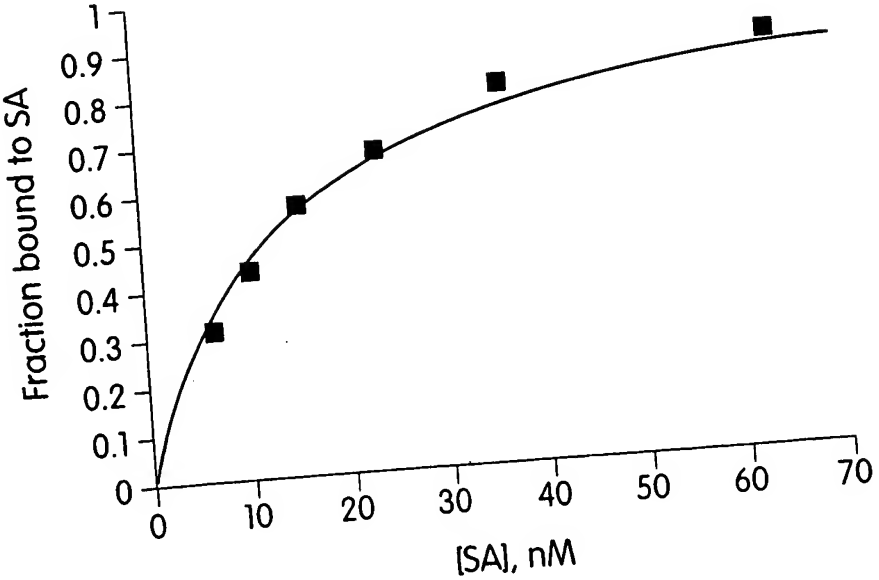

Fig. 4C

		% binding	SEQ ID NO:
FL	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREPLVQEVEDVDEGLVQDLHGVVAGLLDPVEKLLTDWFKKFKNVSKDCKMTFYLEMYDWSGGCKLG	85	21
C1	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREPLVQEVEDVDEGLVQDLHGVVAGLLDPVEKLLTDWFKKFKNVS MMSGGCKLG	87	22
C2	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREPLVQEVEDVDEGLVQDLHGVVAGLLDPVE MMSGGCKLG	88	23
C3	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREPLVQEVEDVDEGLVQ MMSGGCKLG	89	24
C4	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREP MMSGGCKLG	88	25
M1	MDEKTTGWRGGHVVEGLAGELEQLRARLEHHGAGQREP MMSGGCKLG	0.065	26
N1	MD GHVVEGLAGELEQLRARLEHHPQGQREP MMSGGCKLG	69	27
N2	MD EGLAGELEQLRARLEHHPQGQREPLVQEVEDVDEGLVQDLHGVVAGLLDPVEKLLTDWFKKFKNVSKDCKMTFYLEMYDWSGGCKLG	30	28
N3	M ELEQLRARLEHHPQGQREP MMSGGCKLG	0.058	29

Fig. 5

(SEQ ID No.: 37)

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GAGTGCGGCCGCAAGCTTTTAGTCGTCATGTCCATGATAGGTGTCGTCCCCGATATCAATGCTATTGTTAAAGCAGGTCT
TACACATGTTATAGATCCTCAAATGCTTGTTCTTCACCTTCCAGTTCCGGGGAGCCACCTTGCATTTCACACAAGGGTCC
ATGCTACCTCGGGGTACCAAGAATTCGTGATGATGGTGATGGTGACCGGATCCTGGTTCACGTTGACCTTGTGGGTGATG
CTCCAGTCGTGCACGAAGTTGTTCAAGTTCCCCAGCCAGTCCTTCAACAACGTGTCCACCTCGCCAACCAGTGGTCTTCT
CGTCCATCCCTGAGCCGCTACCTCCTGAGCTCGAATTAGTCTGCGCGTCTTTCAGGGCTTCATCGACAGTCTGACGACCG
CTGGCGGCGTTGATCACCGCAGTACGCACGGCATACCAGAAAGCGGACATCTGCGGGATGTTCGGCATGATTTCACCTTT
CTGGGCGTTTTCCATTGTGGCGGCAATACGTGGATCTTTCGCCAACTCTTCCTCGTAAGACTTCAGCGCTACGGCACCCA
GCGGTTTGTCTTTATTAACCGCTTCCAGACCTTCATCAGTCAGCAGATAGTTTTCGAGGAACTCTTTTGCCAGCTCTTTG
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CGTTACACCATAATTCACTTTGCTGGTGTCGATGTTGGACCATGCCCACGGGCCGTTGATGGTCATCGCTGTTTCGCCTT
TATTAAAGGCAGCTTCTGCGATGGAGTAATCGGTGTCTGCATTCATGTGTTTGTTTTTAATCAGGTCAACCAGGAAGGTC
AGACCCGCTTTCGCGCCAGCGTTATCCACGCCCACGTCTTTAATGTCGTACTTGCCGTTTTCATACTTGAACGCATAACC
CCCGTCAGCAGCAATCAGCGGCCAGGTGAAGTACGGTTCTTGCAGGTTGAACATCAGCGCGCTCTTACCTTTCGCTTTCA
GTTCTTTATCCAGCGCCGGGATCTCTTCCCAGGTTTTTGGCGGGTTCGGCAGCAGATCTTTGTTATAAATCAGCGATAAC
GCTTCAACAGCGATCGGGTAAGCAATCAGCTTGCCGTTGTAACGTACGGCATCCCAGGTAAACGGATACAGCTTGTCCTG
GAACGCTTTGTCCGGGGTGATTTCAGCCAACAGGCCAGATTGAGCGTAGCCACCAAAGCGGTCGTGTGCCCAGAAGATAA
TGTCAGGGCCATCGCCAGTTGCCGCAACCTGTGGGAATTTCTCTTCCAGTTTATCCGGATGCTCAACGGTGACTTTAATT
CCGGTATCTTTCTCGAATTTCTTACCGACTTCAGCGAGACCGTTATAGCCTTTATCGCCGTTAATCCAGATTACCAGTTT
ACCTTCTTCGATTCCCATGGTATATCTCCTTCTTAAAGTTAAACAAAATTATTTCTAGAGGGGAATTGTTATCCGCTCAC
AATTCCCCTATAGTGAGTCGTATTAATTTCGCGGGATCGAGATCTCGATCCTCTACGCCGGACGCATCGTGGCCGGCATC
ACCGGCGCCACAGGTGCGGTTGCTGGCGCCTATATCGCCGACATCACCGATGGGGAAGATCGGGCTCGCCACTTCGGGCT
CATGAGCGCTTGTTTCGGCGTGGGTATGGTGGCAGGCCCCGTGGCCGGGGGACTGTTGGGCGCCATCTCCTTGCATGCAC
CATTCCTTGCGGCGGCGGTGCTCAACGGCCTCAACCTACTACTGGGCTGCTTCCTAATGCAGGAGTCGCATAAGGGAGAG
CGTCGAGATCCCGGACACCATCGAATGGCGCAAAACCTTTCGCGGTATGGCATGATAGCGCCCGGAAGAGAGTCAATTCA
GGGTGGTGAATGTGAAACCAGTAACGTTATACGATGTCGCAGAGTATGCCGGTGTCTCTTATCAGACCGTTTCCCGCGTG
GTGAACCAGGCCAGCCACGTTTCTGCGAAAACGCGGGAAAAAGTGGAAGCGGCGATGGCGGAGCTGAATTACATTCCCAA
CCGCGTGGCACAACAACTGGCGGGCAAACAGTCGTTGCTGATTGGCGTTGCCACCTCCAGTCTGGCCCTGCACGCGCCGT

Fig. 6A

GCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCC
GACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCAC
TGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTA
GAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAA
CAAACCACCGCTGGTAGCGGTGGTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCC
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ACTCACCACTGCGATCCCCGGGAAAACAGCATTCCAGGTATTAGAAGAATATCCTGATTCAGGTGAAAATATTGTTGATG
CGCTGGCAGTGTTCCTGCGCCGGTTGCATTCGATTCCTGTTTGTAATTGTCCTTTTAACAGCGATCGCGTATTTCGTCTC
GCTCAGGCGCAATCACGAATGAATAACGGTTTGGTTGATGCGAGTGATTTTGATGACGAGCGTAATGGCTGGCCTGTTGA
ACAAGTCTGGAAAGAAATGCATAAACTTTTGCCATTCTCACCGGATTCAGTCGTCACTCATGGTGATTTCTCACTTGATA
ACCTTATTTTTGACGAGGGGAAATTAATAGGTTGTATTGATGTTGGACGAGTCGGAATCGCAGACCGATACCAGGATCTT
GCCATCCTATGGAACTGCCTCGGTGAGTTTTCTCCTTCATTACAGAAACGGCTTTTTCAAAAATATGGTATTGATAATCC
TGATATGAATAAATTGCAGTTTCATTTGATGCTCGATGAGTTTTTCTAAGAATTAATTCATGAGCGGATACATATTTGAA
TGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAAAGTGCCACCTGAAATTGTAAACGTTAATAT
TTTGTTAAAATTCGCGTTAAATTTTTGTTAAATCAGCTCATTTTTTAACCAATAGGCCGAAATCGGCAAAATCCCTTATA
AATCAAAAGAATAGACCGAGATAGGGTTGAGTGTTGTTCCAGTTTGGAACAAGAGTCCACTATTAAAGAACGTGGACTCC
AACGTCAAAGGGCGAAAAACCGTCTATCAGGGCGATGGCCCACTACGTGAACCATCACCCTAATCAAGTTTTTTGGGGTC
GAGGTGCCGTAAAGCACTAAATCGGAACCCTAAAGGGAGCCCCCGATTTAGAGCTTGACGGGGAAAGCCGGCGAACGTGG
CGAGAAAGGAAGGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGTAGCGGTCACGCTGCGCGTAACCACC
ACACCCGCCGCGCTTAATGCGCCGCTACAGGGCGCGTCCCATTCGCCA

Fig. 6A (continued)

(SEQ ID No.:38)

MGIEEGKLVIWINGDKGYNGLAEVGKKFEKDTGIKVTVEHPDKLEEKFPQVAATGDGPDIIFWAHDRFGGYAQSGLLAEI
TPDKAFQDKLYPFTWDAVRYNGKLIAYPIAVEALSLIYNKDLLPNPPKTWEEIPALDKELKAKGKSALMFNLQEPYFTWP
LIAADGGYAFKYENGKYDIKDVGVDNAGAKAGLTFLVDLIKNKHMNADTDYSIAEAAFNKGETAMTINGPWAWSNIDTSK
VNYGVTVLPTFKGQPSKPFVGVLSAGINAASPNKELAKEFLENYLLTDEGLEAVNKDKPLGAVALKSYEEELAKDPRIAA
TMENAQKGEIMPNIPQMSAFWYAVRTAVINAASGRQTVDEALKDAQTNSSSGGSGSGMDEKTTGWRGGHVVEGLAGELEQ
LRARLEHHPQGQREPGSGHHHHHHEFLVPRGSMDPCVKCKVAPRNWKVKNKHLRIYNMCKTCFNNSIDIGDDTYHGHDD

Fig. 6B

(SEQ ID No.:39)

MDPCVKCKVAPRNWKVKNKHLRIYNMCKTCFNNSIDIGDDTYHGHDD

Fig. 6C

MDEKTTGWRGGHVVEGLAGELEQLRARLEHHPQGQREP

Fig. 7A

Fig. 7B

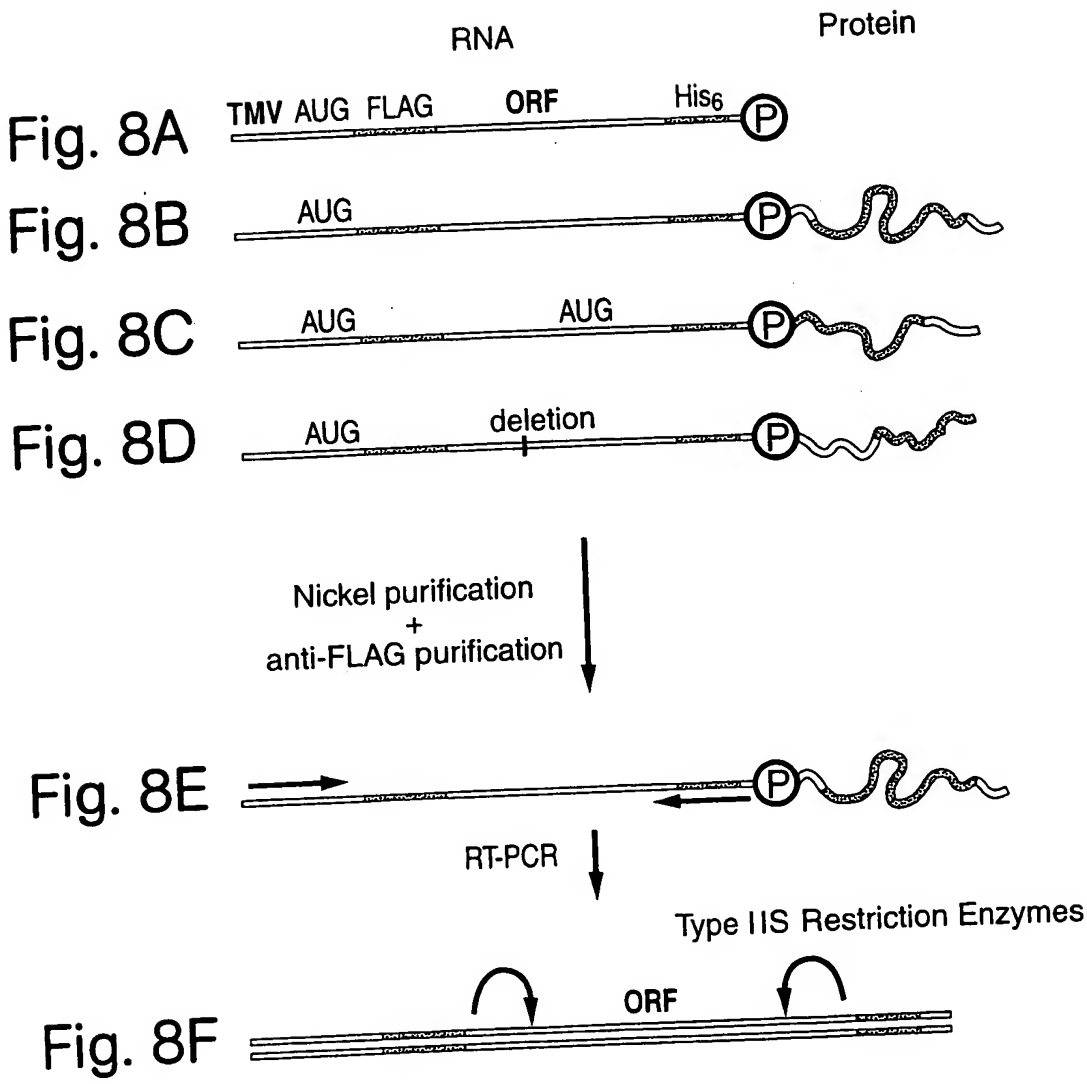

Fig. 8A

Fig. 8B

Fig. 8C

Fig. 8D

Fig. 8E

Fig. 8F

(SEQ ID No.:40)

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GAGTGCGGCCGCAAGCTTCAGCTGCGAAGCTTATTAGAATTCGTGATGATGGTGATGGTGACCGGATCCTGGTTCACGTT
GACCTTGTGGGTGATGCTCCAGTCGTGCACGAAGTTGTTCAAGTTCCCCAGCCAGTCCTTCAACAACGTGTCCACCTCGC
CAACCAGTGGTCTTCTCGTCCATCCCTGAGCCGCTACCTCCTGAGCTCGAATTAGTCTGCGCGTCTTTCAGGGCTTCATC
GACAGTCTGACGACCGCTGGCGGCGTTGATCACCGCAGTACGCACGGCATACCAGAAAGCGGACATCTGCGGGATGTTCG
GCATGATTTCACCTTTCTGGGCGTTTTCCATtGTGGCGGCAATACGTGGATCTTTCGCCAACTCTTCCTCGTAAGACTTC
AGCGCTACGGCACCCAGCGGTTTGTCTTTATTAACCGCTTCCAGACCTTCATCAGTCAGCAGATAGTTTTCGAGGAACTC
TTTTGCCAGCTCTTTGTTCGGACTGGCGGCGTTAATACCTGCGCTCAGCACGCCAACGAACGGTTTGGATGGTTGACCCT
TGAAGGTCGGCAGTACCGTTACACCATAATTCACTTTGCTGGTGTCGATGTTGGACCATGCCCACGGGCCGTTGATGGTC
ATCGCTGTTTCGCCTTTATTAAAGGCAGCTTCTGCGATGGAGTAATCGGTGTCTGCATTCATGTGTTTGTTTTTAATCAG
GTCAACCAGGAAGGTCAGACCCGCTTTCGCGCCAGCGTTATCCACGCCCACGTCTTTAATGTCGTACTTGCCGTTTTCAT
ACTTGAACGCATAACCCCCGTCAGCAGCAATCAGCGGCCAGGTGAAGTACGGTTCTTGCAGGTTGAACATCAGCGCGCTC
TTACCTTTCGCTTTCAGTTCTTTATCCAGCGCCGGGATCTCTTCCCAGGTTTTTGGCGGGTTCGGCAGCAGATCTTTGTT
ATAAATCAGCGATAACGCTTCAACAGCGATCGGGTAAGCAATCAGCTTGCCGTTGTAACGTACGGCATCCCAGGTAAACG
GATACAGCTTGTCCTGGAACGCTTTGTCCGGGGTGATTTCAGCCAACAGGCCAGATTGAGCGTAGCCACCAAAGCGGTCG
TGTGCCCAGAAGATAATGTCAGGGCCATCGCCAGTTGCCGCAACCTGTGGGAATTTCTCTTCCAGTTTATCCGGATGCTC
AACGGTGACTTTAATTCCGGTATCTTTCTCGAATTTCTTACCGACTTCAGCGAGACCGTTATAGCCTTTATCGCCGTTAA
TCCAGATTACCAGTTTACCTTCTTCGATTcccatggTATATCTCCTTCTTAAAGTTAAACAAAATTATTTCTAGAGGGGA
ATTGTTATCCGCTCACAATTCCCCTATAGTGAGTCGTATTAATTTCGCGGGATCGAGATCTCGATCCTCTACGCCGGACG
CATCGTGGCCGGCATCACCGGCGCCACAGGTGCGGTTGCTGGCGCCTATATCGCCGACATCACCGATGGGGAAGATCGGG
CTCGCCACTTCGGGCTCATGAGCGCTTGTTTCGGCGTGGGTATGGTGGCAGGCCCCGTGGCCGGGGGACTGTTGGGCGCC
ATCTCCTTGCATGCACCATTCCTTGCGGCGGCGGTGCTCAACGGCCTCAACCTACTACTGGGCTGCTTCCTAATGCAGGA
GTCGCATAAGGGAGAGCGTCGAGATCCCGGACACCATCGAATGGCGCAAAACCTTTCGCGGTATGGCATGATAGCGCCCG
GAAGAGAGTCAATTCAGGGTGGTGAATGTGAAACCAGTAACGTTATACGATGTCGCAGAGTATGCCGGTGTCTCTTATCA
GACCGTTTCCCGCGTGGTGAACCAGGCCAGCCACGTTTCTGCGAAAACGCGGGAAAAAGTGGAAGCGGCGATGGCGGAGC
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GCCCTGCACGCGCCGTCGCAAATTGTCGCGGCGATTAAATCTCGCGCCGATCAACTGGGTGCCAGCGTGGTGGTGTCGAT
GGTAGAACGAAGCGGCGTCGAAGCCTGTAAAGCGGCGGTGCACAATCTTCTCGCGCAACGCGTCAGTGGGCTGATCATTA

Fig. 9A

ACTATCCGCTGGATGACCAGGATGCCATTGCTGTGGAAGCTGCCTGCACTAATGTTCCGGCGTTATTTCTTGATGTCTCT
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TCACCAGCAAATCGCGCTGTTAGCGGGCCCATTAAGTTCTGTCTCGGCGCGTCTGCGTCTGGCTGGCTGGCATAAATATC
TCACTCGCAATCAAATTCAGCCGATAGCGGAACGGGAAGGCGACTGGAGTGCCATGTCCGGTTTTCAACAAACCATGCAA
ATGCTGAATGAGGGCATCGTTCCCACTGCGATGCTGGTTGCCAACGATCAGATGGCGCTGGGCGCAATGCGCGCCATTAC
CGAGTCCGGGCTGCGCGTTGGTGCGGATATCTCGGTAGTGGGATACGACGATACCGAAGACAGCTCATGTTATATCCCGC
CGTTAACCACCATCAAACAGGATTTTCGCCTGCTGGGGCAAACCAGCGTGGACCGCTTGCTGCAACTCTCTCAGGGCCAG
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TCCCCGCGCGTTGGCCGATTCATTAATGCAGCTGGCACGACAGGTTTCCCGACTGGAAAGCGGGCAGTGAGCGCAACGCA
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CAGCGCTCTGGGTCATTTTCGGCGAGGACCGCTTTCGCTGGAGCGCGACGATGATCGGCCTGTCGCTTGCGGTATTCGGA
ATCTTGCACGCCCTCGCTCAAGCCTTCGTCACTGGTCCCGCCACCAAACGTTTCGGCGAGAAGCAGGCCATTATCGCCGG
CATGGCGGCCCCACGGGTGCGCATGATCGTGCTCCTGTCGTTGAGGACCCGGCTAGGCTGGCGGGGTTGCCTTACTGGTT
AGCAGAATGAATCACCGATACGCGAGCGAACGTGAAGCGACTGCTGCTGCAAAACGTCTGCGACCTGAGCAACAACATGA
ATGGTCTTCGGTTTCCGTGTTTCGTAAAGTCTGGAAACGCGGAAGTCAGCGCCCTGCACCATTATGTTCCGGATCTGCAT
CGCAGGATGCTGCTGGCTACCCTGTGGAACACCTACATCTGTATTAACGAAGCGCTGGCATTGACCCTGAGTGATTTTTC
TCTGGTCCCGCCGCATCCATACCGCCAGTTGTTTACCCTCACAACGTTCCAGTAACCGGGCATGTTCATCATCAGTAACC
CGTATCGTGAGCATCCTCTCTCGTTTCATCGGTATCATTACCCCCATGAACAGAAATCCCCCTTACACGGAGGCATCAGT
GACCAAACAGGAAAAAACCGCCCTTAACATGGCCCGCTTTATCAGAAGCCAGACATTAACGCTTCTGGAGAAACTCAACG
AGCTGGACGCGGATGAACAGGCAGACATCTGTGAATCGCTTCACGACCACGCTGATGAGCTTTACCGCAGCTGCCTCGCG
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GCGGAGTGTATACTGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGCACCATATATGCGGTGTGAAATACC
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ATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCC
TGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCCC
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GTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCGCTCCAAGCTGGGCTGTGTGCACGA
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Fig. 9A (continued)

CACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCT
AACTACGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAG
CTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAAG
GATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTC
ATGAACAATAAAACTGTCTGCTTACATAAACAGTAATACAAGGGGTGTTATGAGCCATATTCAACGGGAAACGTCTTGCT
CTAGGCCGCGATTAAATTCCAACATGGATGCTGATTTATATGGGTATAAATGGGCTCGCGATAATGTCGGGCAATCAGGT
GCGACAATCTATCGATTGTATGGGAAGCCCGATGCGCCAGAGTTGTTTCTGAAACATGGCAAAGGTAGCGTTGCCAATGA
TGTTACAGATGAGATGGTCAGACTAAACTGGCTGACGGAATTTATGCCTCTTCCGACCATCAAGCATTTTATCCGTACTC
CTGATGATGCATGGTTACTCACCACTGCGATCCCCGGGAAAACAGCATTCCAGGTATTAGAAGAATATCCTGATTCAGGT
GAAAATATTGTTGATGCGCTGGCAGTGTTCCTGCGCCGGTTGCATTCGATTCCTGTTTGTAATTGTCCTTTTAACAGCGA
TCGCGTATTTCGTCTCGCTCAGGCGCAATCACGAATGAATAACGGTTTGGTTGATGCGAGTGATTTTGATGACGAGCGTA
ATGGCTGGCCTGTTGAACAAGTCTGGAAAGAAATGCATAAACTTTTGCCATTCTCACCGGATTCAGTCGTCACTCATGGT
GATTTCTCACTTGATAACCTTATTTTTGACGAGGGGAAATTAATAGGTTGTATTGATGTTGGACGAGTCGGAATCGCAGA
CCGATACCAGGATCTTGCCATCCTATGGAACTGCCTCGGTGAGTTTTCTCCTTCATTACAGAAACGGCTTTTTCAAAAAT
ATGGTATTGATAATCCTGATATGAATAAATTGCAGTTTCATTTGATGCTCGATGAGTTTTTCTAAGAATTAATTCATGAG
CGGATACATATTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAAAGTGCCACCTGAAA
TTGTAAACGTTAATATTTTGTTAAAATTCGCGTTAAATTTTTGTTAAATCAGCTCATTTTTTAACCAATAGGCCGAAATC
GGCAAAATCCCTTATAAATCAAAAGAATAGACCGAGATAGGGTTGAGTGTTGTTCCAGTTTGGAACAAGAGTCCACTATT
AAAGAACGTGGACTCCAACGTCAAAGGGCGAAAAACCGTCTATCAGGGCGATGGCCCACTACGTGAACCATCACCCTAAT
CAAGTTTTTTGGGGTCGAGGTGCCGTAAAGCACTAAATCGGAACCCTAAAGGGAGCCCCCGATTTAGAGCTTGACGGGGA
AAGCCGGCGAACGTGGCGAGAAAGGAAGGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGTAGCGGTCAC
GCTGCGCGTAACCACCACACCCGCCGCGCTTAATGCGCCGCTACAGGGCGCGTCCCATTCGCCA

Fig. 9A (continued)

(SEQ ID No.:41)

MGIEEGKLVIWINGDKGYNGLAEVGKKFEKDTGIKVTVEHPDKLEEKFPQVAATGDGPDIIFWAHDRFGGYAQSGLLAEI
TPDKAFQDKLYPFTWDAVRYNGKLIAYPIAVEALSLIYNKDLLPNPPKTWEEIPALDKELKAKGKSALMFNLQEPYFTWP
LIAADGGYAFKYENGKYDIKDVGVDNAGAKAGLTFLVDLIKNKHMNADTDYSIAEAAFNKGETAMTINGPWAWSNIDTSK
VNYGVTVLPTFKGQPSKPFVGVLSAGINAASPNKELAKEFLENYLLTDEGLEAVNKDKPLGAVALKSYEEELAKDPRIAA
TMENAQKGEIMPNIPQMSAFWYAVRTAVINAASGRQTVDEALKDAQTNSSSGGSGSGMDEKTTGWRGGHVVEGLAGELEQ
LRARLEHHPQGQREPGSGHHHHHHEF

Fig. 9B

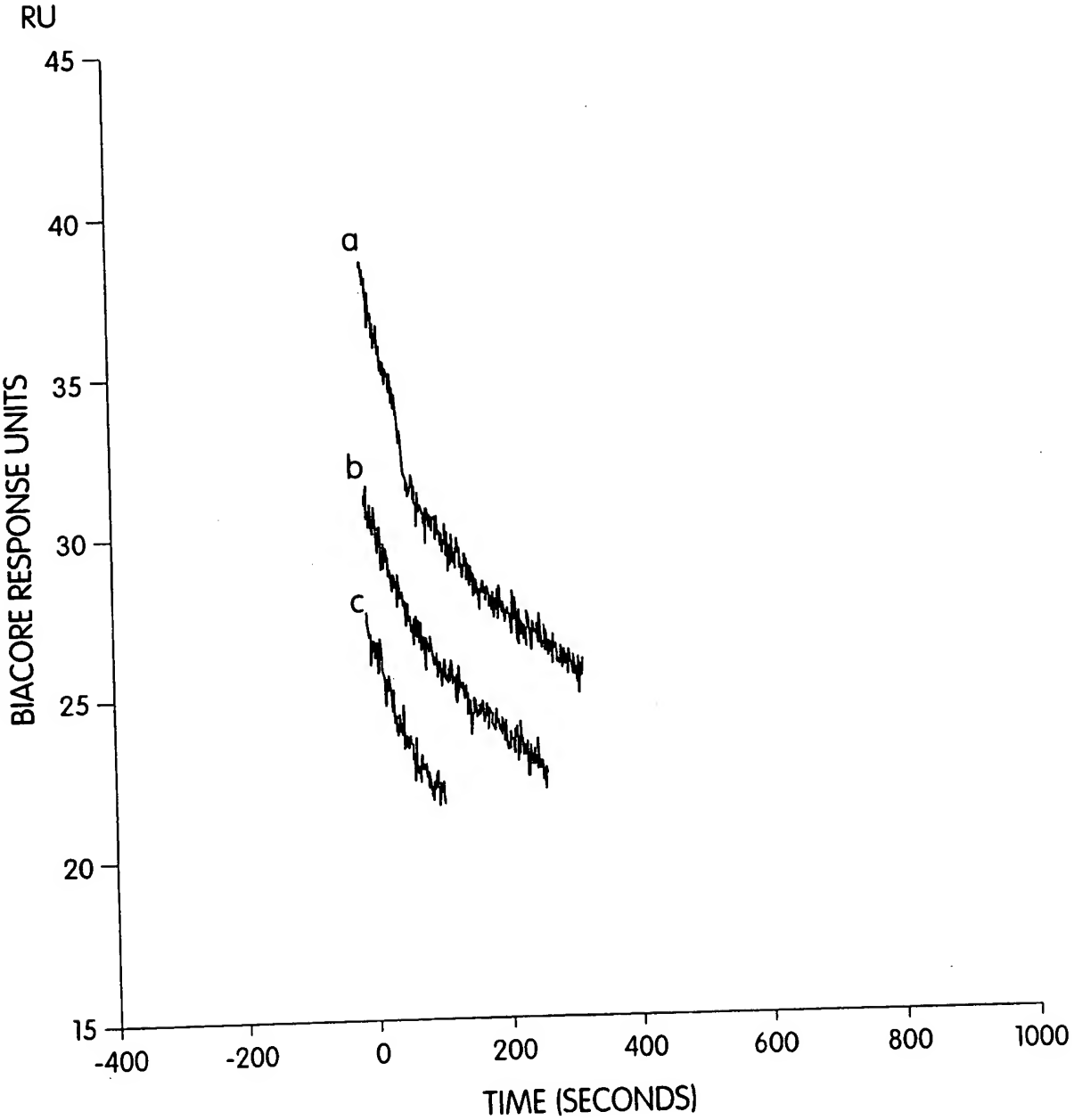

Fig. 10A

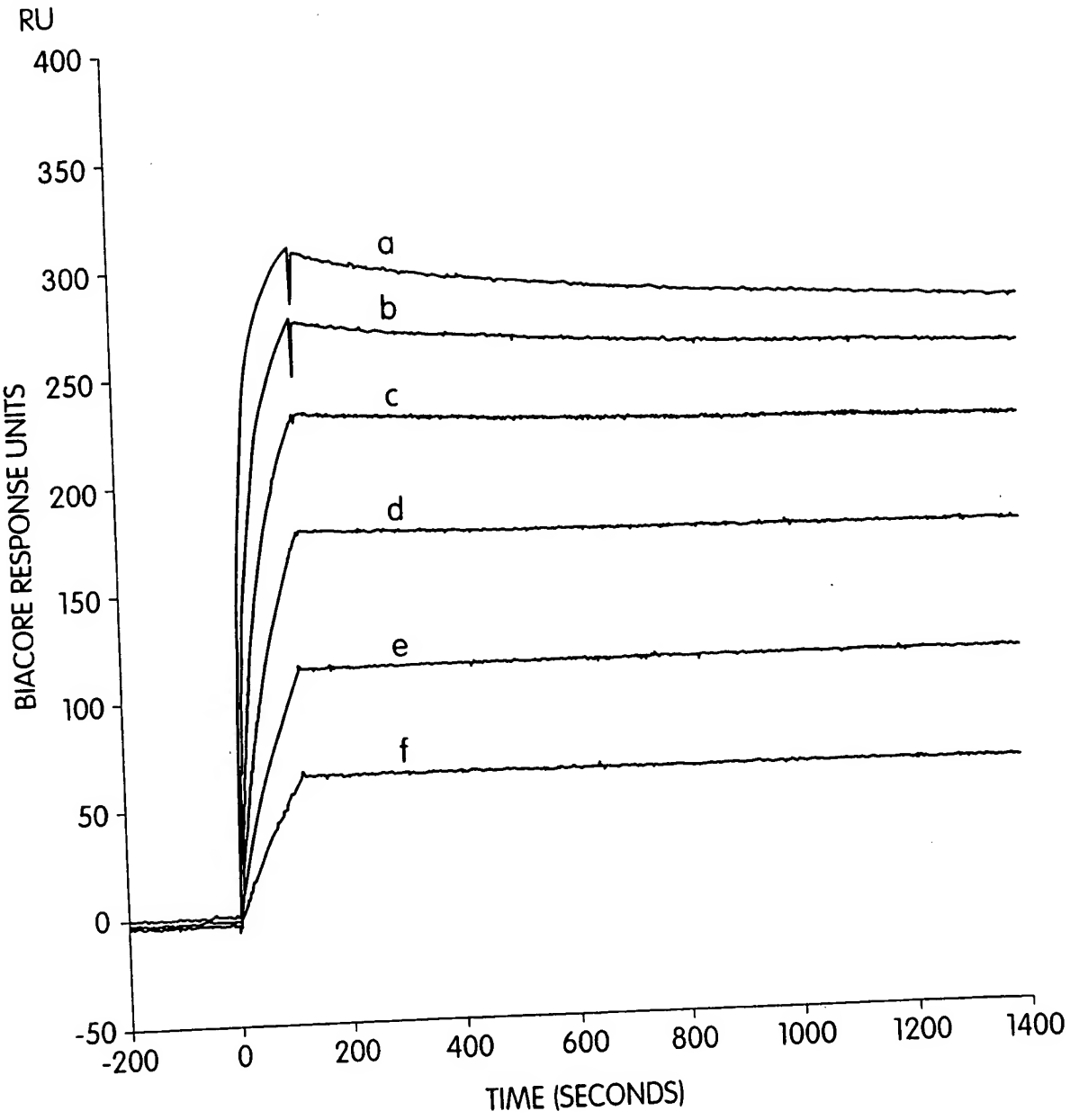
RU
400
350
300
250
200
150
100
50
0
-50
BIACORE RESPONSE UNITS
a
b
c
d
e
f
-200
0
200
400
600
800
1000
1200
1400
TIME (SECONDS)

Fig. 10B

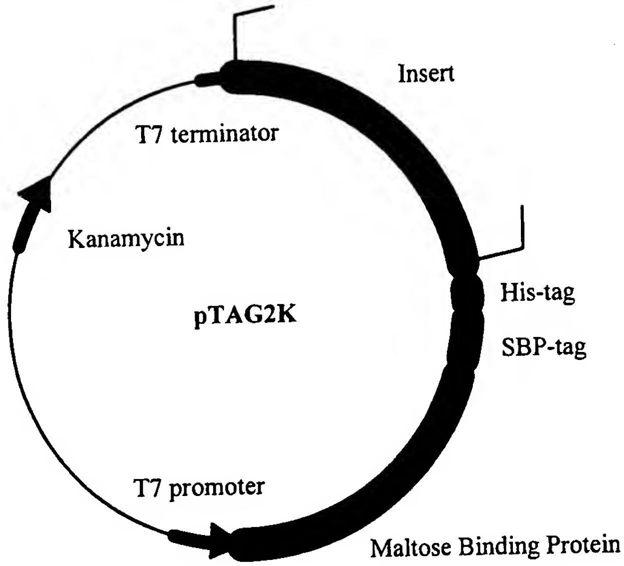

Fig. 11

Proportion of binding inhibited
1.1
0.9
0.7
0.5
0.3
0.1
-0.1
0.01
0.1
1
10
100
1000
10000
[Streptavidin], nM

Fig. 12

Uninduced
Induced
Soluble Fraction
Biotin Elution
Imidazole Elution
Maltose Elution
83 kDa >
62 kDa >
47.5 kDa >
32.5 kDa >
25 kDa >

Fig. 13

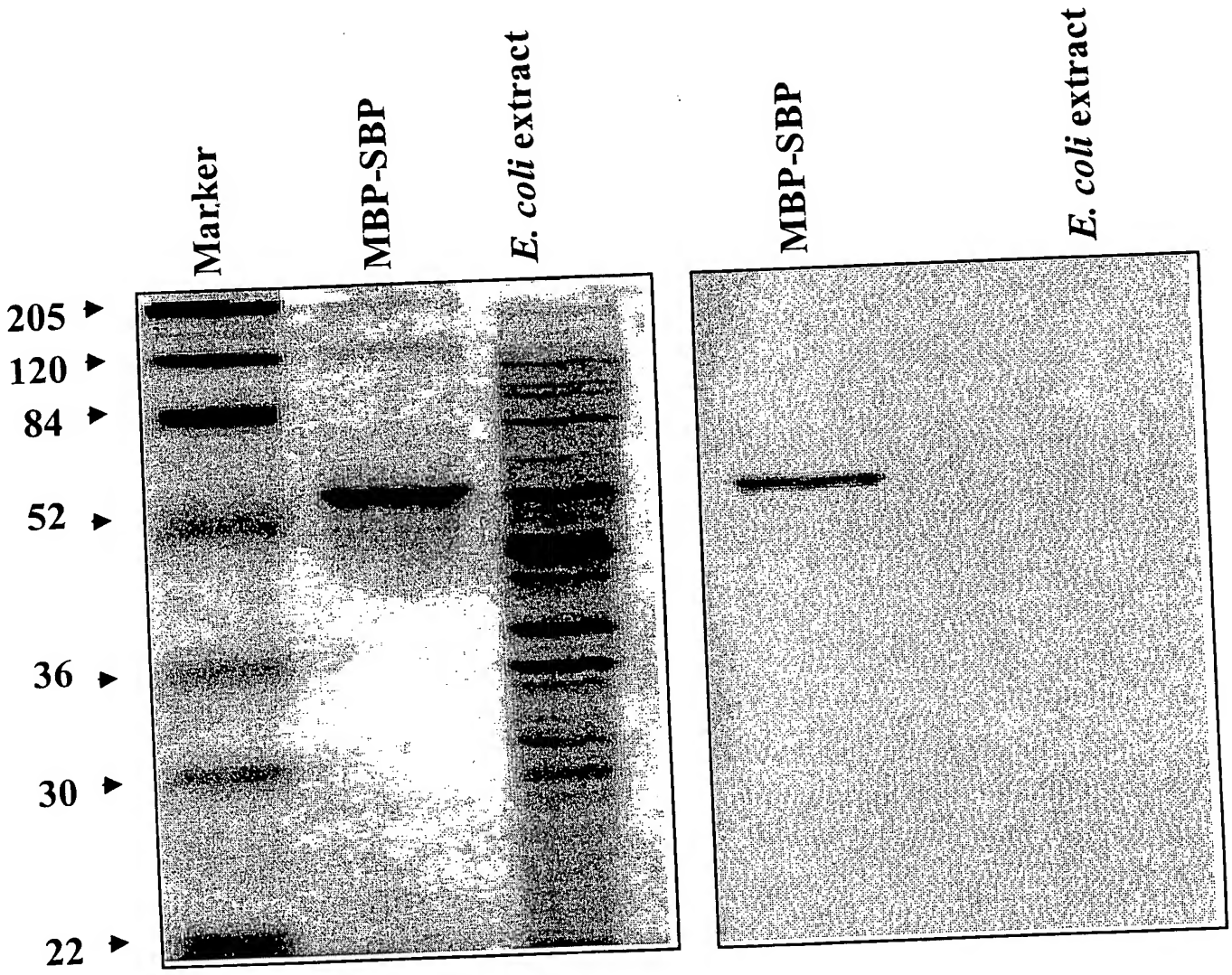

Fig. 14

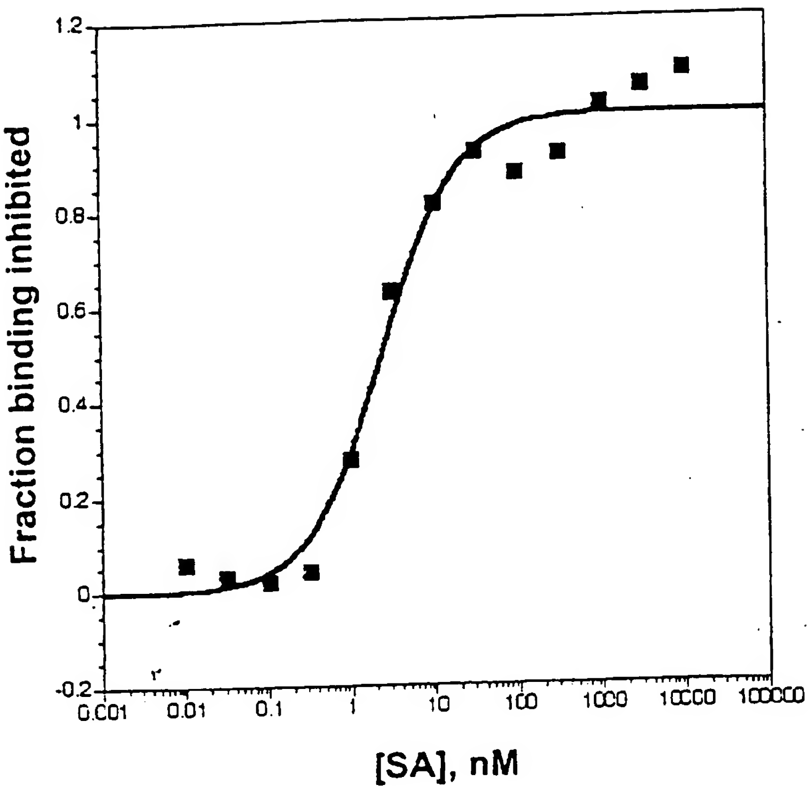
1.2
1
0.8
0.6
0.4
0.2
0
-0.2
Fraction binding inhibited
0.001
0.01
0.1
1
10
100
1000
10000
100000
[SA], nM

Fig. 15